

FORM PTO-1449 (Modified)

ATTY. DOCKET NO.
24737-1906B

SERIAL NO.
09/704,362

LIST OF PATENTS AND PUBLICATIONS FOR
APPLICANT'S INFORMATION DISCLOSURE
STATEMENT

APPLICANT
Ramnarayan *et al.*

FILING DATE
November 1, 2000

GROUP
1631



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U.S. PATENT DOCUMENTS

EXAMINER INITIAL		DOCUMENT NUMBER							DATE	NAME	CLASS	SUB CLASS	FILING DATE
J83	AA	5	3	1	7	0	9	7	05/31/94	Miller <i>et al.</i>	536	24.31	10/07/91
J83	AB	5	4	9	5	4	2	3	02/27/96	DeLisi <i>et al.</i>	364	496	10/25/93
J83	AC	5	5	9	3	9	5	9	01/14/97	Miller <i>et al.</i>	514	8	10/14/93
J83	AD	5	6	2	4	8	1	7	04/29/97	Miller <i>et al.</i>	435	69.1	04/28/94
J83	AE	5	6	9	9	2	6	8	12/16/97	Schmidt	364	496	06/07/95
J83	AF	5	9	6	8	7	3	7	10/19/99	Ali-Osman <i>et al.</i>	435	6	11/12/96
J83	AG	5	9	7	8	7	4	0	11/02/99	Armistead <i>et al.</i>	702	19	08/09/95
J83	AH	6	1	2	8	5	8	2	10/03/00	Wilson <i>et al.</i>	702	27	04/30/96

FOREIGN PATENT DOCUMENTS

		DOCUMENT NUMBER							DATE	COUNTRY	CLASS	SUB CLASS	Translation Yes No	
None														

OTHER ART (Including Author, Title, Date, Pertinent Pages, Etc.)

J83	AI	Baker <i>et al.</i> , "Protein Structure Prediction and Structural Genomics", Science, <u>294</u> :93-96 (2001)											
J83	AJ	Hess <i>et al.</i> , "Impact of Pharmacogenomics on the Clinical Laboratory", Mol. Diagn., <u>4(4)</u> :289-98 (1999)											
J83	AK	Hess <i>et al.</i> , "Gene Therapy Monitoring: Clinical Monitoring for Efficacy and Potential Toxicity, Mol. Diagn., <u>2(2)</u> :147-155 (1997)											

EXAMINER

J83. Bence

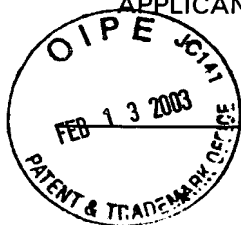
DATE CONSIDERED

4/1/03

EXAMINER: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

Title: **USE OF COMPUTATIONALLY DERIVED PROTEIN STRUCTURES OF GENETIC POLYMORPHISMS IN PHARMACOGENOMICS FOR DRUG DESIGN AND CLINICAL APPLICATIONS**

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U.S. PATENT DOCUMENTS

EXAMINER INITIAL		DOCUMENT NUMBER							DATE	NAME	CLASS	SUB CLASS	FILING DATE
JLB	AA	5	7	3	6	5	0	9	04/07/98	Balaji <i>et al.</i>			

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None													

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OTHER ART (Including Author, Title, Date, Pertinent Pages, Etc.)

JLB	AB	Eisenhaber <i>et al.</i> , "Protein structure prediction: recognition of primary, secondary, and tertiary structural features from amino acid sequence", <i>Crit. Rev. Biochem. Mol. Biol.</i> , 30:1-94, 1995
JLB	AC	de Dios <i>et al.</i> , "Secondary and Tertiary Structural Effects on Protein NMR Chemical Shifts: An <i>ab Initio</i> Approach", <i>Science</i> , 260:1491-1496, 1993
JLB	AD	Dunbrack <i>et al.</i> , "Meeting review: the Second Meeting on the Critical Assessment of Techniques for Protein Structure Prediction (CASP2), Asilomar, California, Dec. 13-16, 1996", <i>Folding and Design</i> , 1997, pp. R27-R42
JLB	AE	Jones, "Successful <i>ab initio</i> prediction of the tertiary structure of NK-Lysin using multiple sequences and recognized supersecondary structural motifs", <i>Proteins: Structure, function, and Genetics</i> , Supp. 1, 1997, pp. 185-191
JLB	AF	Osguthorpe, "Improved <i>Ab Initio</i> Predictions with a Simplified, Flexible Geometry Model", <i>Proteins: Structure, Function, and Genetics</i> , Suppl. 3 (November 9, 1999), pp. 186-193
JLB	AG	Samudrala <i>et al.</i> , "Ab initio protein structure prediction using a combined hierarchical approach", <i>Proteins: Structure, Function, and Genetics</i> , Supp. 3, 194-198, 1999
JLB	AH	Westhead and Thornton, "Protein structure prediction", <i>Curr. Opin. Biotechnol.</i> , 9:383-389, 1998

EXAMINER

JLB. Buser

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